

AMENDMENTS TO THE CLAIMS

Please amend the claims as follows:

1. (currently amended) A focused library of vectors or genetic packages that display, display and express, or comprise a member of a diverse family of human antibody related peptides, polypeptides and proteins and collectively display, display and express, or comprise at least a portion of the diversity of the antibody family, the vectors or genetic packages being characterized by variegated DNA sequences that encode a heavy chain CDR1 selected from the group consisting of:

(1) $\langle 1 \rangle_1 Y_2 \langle 1 \rangle_3 M_4 \langle 1 \rangle_5$, wherein $\langle 1 \rangle$ is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y;

(2) $(S/T)_1 (S/G/X)_2 (S/G/X)_3 Y_4 Y_5 W_6 (S/G/X)_7$. wherein (S/T) is a 1:1 mixture of S and T residues, (S/G/X) is a mixture of 0.2025 S, 0.2025 G and 0.035 of each of amino acid residues A, D, E, F, H, I, K, L, M, N, P, Q, R, T, V, W, and Y;

(3) $V_1 S_2 G_3 G_4 S_5 I_6 S_7 \langle 1 \rangle_8 \langle 1 \rangle_9 \langle 1 \rangle_{10} Y_{11} Y_{12} W_{13} \langle 1 \rangle_{14}$ (**SEQ ID NO: 1**), wherein $\langle 1 \rangle$ is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y; and

(4) mixtures of vectors or genetic packages characterized by any of the above DNA sequences.

2. (original) The focused library according to claim 1, wherein HC CDR1s (1), (2) and (3) are present in the library in the ratio 0.80:0.17:0.02.

3. (currently amended) A focused library of vectors or genetic packages that display, display and express, or comprise a member of a diverse family of human antibody related peptides, polypeptides and proteins and collectively display, display and express, or comprise at least a portion of the diversity of the antibody facility, the vectors or genetic packages being characterized by variegated DNA sequences that encode a heavy chain CDR2 selected from the group consisting of:

(1) <2>I<2><3>SGG<1>T<1>YADSVKG (SEQ ID NO: 2), wherein <1> is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y; <2> is an equimolar mixture of each of amino acid residues Y, R, W, V, G, and S; and <3> is an equimolar mixture of each of amino acid residues P, S, and G or an equimolar mixture of P and S;

(2) <1>I<4><1><1><G><5><1><1><1>YADSVKG (SEQ ID NO: 3), wherein <1> is an equimolar mixture of each of

amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y; <4> is an equimolar mixture of residues D, I, N, S, W, Y; and <5> is an equimolar mixture of residues S, G, D and N;

(3) <1>I<4><1><1>G<5><1><1>YNPSLK (SEQ ID NO: 4), wherein <1> is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and <4> and <5> are as defined above;

(4) <1>I<8>S<1><1><1>GGYY<1>YAASVK (SEQ ID NO: 5), wherein <1> is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; <8> is 0.27 R and 0.027 of each of ADEFGHIKLMNPQSTVWY; and

(5) mixtures of vectors or genetic packages characterized by any of the above DNA sequences.

4. (original) The focused library according to claim 3 wherein a mixture of HC CDR2s (1)/(2) (equimolar), (3) and (4) are present in the library in a ratio of 0.54:0.43:0.03.

5. (currently amended) A focused library of vectors or genetic packages that display, display and express, or comprise a member of a diverse family of human antibody related peptides, polypeptides and

proteins and collectively display, display and express,
or comprise at least a portion of the diversity of the
antibody family, the vectors or genetic packages being
characterized by variegated DNA sequences that encode a
heavy chain CDR3 selected from the group consisting of:

(1) YYCA21111YFDYWG (SEQ ID NO: 6), wherein 1
is an equimolar mixture of each amino acid residues A, D,
E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and
2 is an equimolar mixture of K and R;

(2) YYCA211111YFDYWG (SEQ ID NO: 7), wherein
1 is an equimolar mixture of each amino acid residues A,
D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y;
and 2 is an equimolar mixture of K and R;

(3) YYCA21111111YFDYWG (SEQ ID NO: 8),
wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; and 2 is an equimolar mixture of K and R;

(4) YYCAR111S2S3111YFDYWG (SEQ ID NO: 9),
wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; and 2 is an equimolar mixture of S and G; and
3 is an equimolar mixture of Y and W;

(5) YYCA2111CSG11CY1YFDYWG (SEQ ID NO: 10),
wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; and 2 is an equimolar mixture of K and R;

(6) YYCA2111S1TIFG11111YFDYWG (SEQ ID NO: 11),
wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; and 2 is an equimolar mixture of K and R;

(7) YYCAR111YY2S3344111YFDYWG (SEQ ID NO: 12),
wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; 2 is an equimolar mixture of D and S; and 3
is an equimolar mixture of S and G;

(8) YYCAR1111YC2231CY111YFDYWG (SEQ ID NO:
13), wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; 2 is an equimolar mixture of S and G; and 3
is an equimolar mixture of T, D and G; and

(9) mixtures of vectors or genetic packages
characterized by any of the above DNA sequences.

6. (original) The focused library according to
claim 5, wherein 1 in one or all of HC CDR3s (1) through

(8) is 0.095 of each of G and Y and 0.048 of each of A, D, E, F, H, I, K, L, M, N, P, Q, R, S, T, V, and W.

7. (original) The focused library according to claim 5 or 6, wherein HC CDR3s (1) through (8) are present in the library in the following proportions:

- (1) 0.10
- (2) 0.14
- (3) 0.25
- (4) 0.13
- (5) 0.13
- (6) 0.11
- (7) 0.04 and
- (8) 0.10

8. (original) The focused library according to claim 5 or 6, wherein the HC CDR3s (1) through (8) are present in the library in the following proportions:

- (1) 0.02
- (2) 0.14
- (3) 0.25
- (4) 0.14
- (5) 0.14
- (6) 0.12
- (7) 0.08 and
- (8) 0.11

9. (currently amended) A focused library of vectors or genetic packages that display, display and express, or comprise a member of a diverse family of

human antibody related peptides, polypeptides and proteins and collectively display, display and express, or comprise at least a portion of the diversity of the antibody family, the vectors or genetic packages being characterized by variegated DNA sequences that encodes a kappa light chain CDR1 selected from the group consisting of:

(1) RASQ<1>V<2><2><3>LA (SEQ ID NO: 14)

(2) RASQ<1>V<2><2><2><3>LA (SEQ ID NO: 15);

wherein <1> is an equimolar mixture of amino acid residues ADEFGHIKLMNPQRSTVWY; <2> is 0.2 S and 0.044 of each of ADEFGHIKLMNPQRTVWY; and <3> is 0.2Y and 0.044 each of ADEFGHIKLMNPQRTVW and Y; and

(3) mixtures of vectors or genetic packages characterized by any of the above DNA sequences.

10. (original) The focused library of claim 9, wherein CDR1s (1) and (2) are present in the library in a ratio of 0.68:0.32.

11. (original) A focused library of vectors or genetic packages that display, display and express, or comprise a member of a diverse family of human antibody related peptides, polypeptides and proteins and collectively display, display and express, or comprise at

least a portion of the diversity of the antibody family,
the vectors or genetic packages being characterized by
variegated DNA sequences that encode a kappa light chain
CDR2 having the sequence:

<1>AS<2>R<4><1>,

wherein <1> is an equimolar mixture of amino acid
residues ADEFGHIKLMNPQRSTVWY; <2> is 0.2 S and 0.044 of
each of ADEFGHIKLMNPQRTVWY; and <4> is 0.2 A and)0.044
each of DEFGHIKLMNPQRSTVWY.

12. (currently amended) A focused library of
vectors or genetic packages that display, display and
express, or comprise a member of a diverse family of
human antibody related peptides, polypeptides and
proteins and collectively display, display and express,
or comprise at least a portion of the diversity of the
antibody family, the vectors or genetic packages being
characterized by variegated DNA sequences that encode a
kappa light chain CDR3 selected from the groups
consisting of:

(1) QQ<3><1><1><1>P<1>T (SEQ ID NO: 16),

wherein <1> is an equimolar mixture of amino
acid residues ADEFGHIKLMNPQRSTVWY; <3> is 0.2 Y and 0.044
each of ADEFGHIKLMNPQRTVW;

(2) QQ33111P, wherein 1 and 3 are as defined
in (1) above;

(3) QQ3211PP1T (SEQ ID NO: 17), wherein 1 and
3 are as defined in (1) above and 2 is 0.2 S and 0.044
each of ADEFGHIKLMNPQRTVWY; and

(4) mixtures of vectors or genetic packages
characterized by any of the above DNA sequences.

13. (original) The focused library according to
claim 12, wherein CDR3s (1), (2) and (3) are present in
the library in a ratio of 0.65:0.1:0.25.

14. (currently amended) A focused library of
vectors or genetic packages that display, display and
express, or comprise a member of a diverse family of
human antibody related peptides, polypeptides and
proteins and collectively display, display and express,
or comprise at least a portion of the diversity of the
antibody family, the vectors or genetic packages being
characterized by variegated DNA sequences that encode a
lambda light chain CDR1 selected from the group
consisting of:

(1) TG<1>SS<2>VG<1><3><2><3>VS (SEQ ID NO:
18),

wherein <1> is 0.27 T, 0.27 G and 0.027 each of ADEFHIKLMNPQRSVWY, <2> is 0.27 D, 0.27 N and 0.027 each of AEFGHIKLMPQRSTVWY, and <3> is 0.36 Y and 0.036 each of ADEFGHIKLMNPQRSTVW;

(2) G<2><4>L<4><4><4><3><4><4>,

wherein <2> is as defined in (1) above and <4> is an equimolar mixture of amino acid residues ADEFGHIKLMNPQRSTVWY; and

(3) mixtures of vectors or genetic packages characterized by any of the above DNA sequences.

15. (original) The focused library according to claim 14, where CDR1s (1) and (2) are present in the library in a ratio of 0.67:0.33.

16. (original) A focused library of vectors or genetic packages that display, display and express, or comprise a member of a diverse family of human antibody related peptides, polypeptides and proteins and collectively display, display and express, or comprise at least a portion of the diversity of the antibody family, the vectors or genetic packages being characterized by variegated DNA sequences that encode a lambda light chain CDR2 has the sequence:

<4><4><4><2>RPS,

wherein <2> is 0.27 D, 0.27 N, and 0.027 each of
AEFGHIKLMPQIRSTVWY and <4> is an equimolar mixture of
amino acid residues ADEFGHIKLMNPQIRSTVW.

17. (currently amended) A focused library of
vectors or genetic packages that display, display and
express, or comprise a member of a diverse family of
human antibody related peptides, polypeptides and
proteins and collectively display, display and express,
or comprise at least a portion of the diversity of the
antibody family, the vectors or genetic packages being
characterized by variegated DNA sequences that encode a
lambda light chain CDR3 selected from the group
consisting of:

(1) <4><5><4><2><4>S<4><4><4><4>V,

wherein <2> is 0.27 D, 0.27 N, and 0.027 each
of AEFGHIKLMPQIRSTVWY; <4> is an equimolar mixture of
amino acid residues ADEFGHIKLMNPQIRSTVW; and <5> is 0.36 S
and 0.0355 each of ADEFGHIKLMNPQRTVWY;

(2) <5>SY<1><5>S<5><1><4>V (SEQ ID NO: 19),

wherein <1> is an equimolar mixture of
ADEFGHIKLMNPQIRSTVWY; and <4> and <5> are as defined in
(1) above; and

(3) mixtures of vectors or genetic packages
characterized by any of the above DNA sequences.

18. (original) The focused library according to claim 17, wherein CDR3s (1) and (2) are present in the library in an equimolar mixture.

19. (original) The focused library according to claim 1 or 2 further comprising variegated DNA sequences that encode a heavy chain CDR selected from the group consisting of:

(1) one or more of the heavy chain CDR2s defined in claim 3 or 4;

(2) one or more of the heavy chain CDR3s defined in claims 5, 6, 7, or 8; and

(3) mixtures of vectors or genetic packages characterized by (1) and (2).

20. (original) The focused library according to claim 3 further comprising variegated DNA sequences that encodes one or more heavy chain CDR3s selected from the group defined in claims 5, 6, 7 or 8.

21. (original) The focused library according to claim 19 or 20, further comprising variegated DNA sequences that encodes a light chain CDR selected from the group consisting of

(1) one or more the kappa light chain CDR1s defined in claim 9 or 10;

- (2) the kappa light chain CDR2 defined in claim 11;
- (3) one or more of the kappa light chain CDR3s defined in claim 12 or 13;
- (4) one or more of the kappa light chain CDR1s defined in claim 14 or 15;
- (5) the lambda light chain CDR2 defined in claim 16;
- (6) one or more of the lambda light chain CDR3s defined in claim 17 or 18; and
- (7) mixtures of vectors and genetic packages characterized by one or more of (1) through (6).

22. (currently amended) A population of variegated DNA sequences that encode a heavy chain CDR1 selected from the group consisting of:

- (1) $\langle 1 \rangle_1 Y_2 \langle 1 \rangle_3 M_4 \langle 1 \rangle_5$, wherein $\langle 1 \rangle$ is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y;
- (2) $(S/T)_1 (S/G/X)_2 (S/G/X)_3 Y_4 Y_5 W_6 (S/G/X)_7$. wherein (S/T) is a 1:1 mixture of S and T residues, (S/G/X) is a mixture of 0.2025 S, 0.2025 G and 0.035 of each of amino acid residues A, D, E, F, H, I, K, L, M, N, P, Q, R, T, V, W, and Y;

(3) $V_1S_2G_3G_4S_5I_6S_7<1>_8<1>_9<1>_{10}Y_{11}Y_{12}W_{13}<1>_{14}$ (SEQ ID NO: 1), wherein $<1>$ is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y; and

(4) mixtures of variegated DNA sequences characterized by any of the above DNA sequences.

23. (original) The population of variegated DNA sequences according to claim 22, wherein HC CDR1s (1), (2) and (3) are present in the population in the ratio 0.80:0.17:0.02.

24. (currently amended) A population of variegated DNA sequences that encode a heavy chain CDR2 selected from the group consisting of:

(1) $<2>I<2><3>SGG<1>T<1>YADSVKG$ (SEQ ID NO: 2), wherein $<1>$ is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y; $<2>$ is an equimolar mixture of each of amino acid residues Y, R, W, V, G, and S; and $<3>$ is an equimolar mixture of each of amino acid residues P, S, and G or an equimolar mixture of P and S;

(2) $<1>I<4><1><1><G><5><1><1><1>YADSVKG$ (SEQ ID NO: 3), wherein $<1>$ is an equimolar mixture of each of

amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, and Y; <4> is an equimolar mixture of residues D, I, N, S, W, Y; and <5> is an equimolar mixture of residues S, G, D and N;

(3) <1>I<4><1><1>G<5><1><1>YNPSLKG (SEQ ID NO: 4), wherein <1> is an equimolar mixture of each of amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and <4> and <5> are as defined above;

(4) <1>I<8>S<1><1><1>GGYY<1>YAASVKG (SEQ ID NO: 5), wherein <1> is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; <8> is 0.27 R and 0.027 of each of ADEFGHIKLMNPQSTVWY; and

(5) mixtures of variegated DNA sequences characterized by any of the above DNA sequences.

25. (original) The population of variegated DNA sequences according to claim 24, wherein a mixture of HC CDR2s (1)/(2) (equimolar), (3) and (4) are present in the population in a ratio of 0.54:0.43:0.03.

26. (currently amended) A population of variegated DNA sequences that encode a heavy chain CDR3 selected from the group consisting of:

(1) YYCA21111YFDYWG (SEQ ID NO: 6), wherein 1 is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and 2 is an equimolar mixture of K and R;

(2) YYCA2111111YFDYWG (SEQ ID NO: 7), wherein 1 is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and 2 is an equimolar mixture of K and R;

(3) YYCA211111111YFDYWG (SEQ ID NO: 8), wherein 1 is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and 2 is an equimolar mixture of K and R;

(4) YYCAR111S2S3111YFDYWG (SEQ ID NO: 9), wherein 1 is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and 2 is an equimolar mixture of S and G; and 3 is an equimolar mixture of Y and W;

(5) YYCA2111CSG11CY1YFDYWG (SEQ ID NO: 10), wherein 1 is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and 2 is an equimolar mixture of K and R;

(6) YYCA211S1TIFG11111YFDYWG (SEQ ID NO: 11), wherein 1 is an equimolar mixture of each amino acid residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W and Y; and 2 is an equimolar mixture of K and R;

(7) YYCAR111YY2S3344111YFDYWG (SEQ ID NO: 12),
wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; 2 is an equimolar mixture of D and S; and 3
is an equimolar mixture of S and G;

(8) YYCAR1111YC2231CY111YFDYWG (SEQ ID NO:
13), wherein 1 is an equimolar mixture of each amino acid
residues A, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T,
V, W and Y; 2 is an equimolar mixture of S and G; and 3
is an equimolar mixture of T, D and G; and

(9) mixtures of variegated DNA sequences
characterized by any of the above DNA sequences.

27. (original) The population of variegated DNA
according to claim 26, wherein 1 in one or all of HC
CDR3s (1) through (8) is 0.095 of each of G and Y and
0.048 of each of A, D, E, F, H, I, K, L, M, N, P, Q, R,
S, T, V, and W.

28. (original) The population of variegated DNA
sequences according to claim 26 or 27, wherein HC CDR3s
(1) through (8) are present in the population in the
following proportions:

- (1) 0.10
- (2) 0.14
- (3) 0.25
- (4) 0.13

- (5) 0.13
- (6) 0.11
- (7) 0.04 and
- (8) 0.10

29. (original) The population of variegated DNA sequences according to claim 26 or 27, wherein the HC CDR3s (1) through (8) are present in the population in the following proportions:

- (1) 0.02
- (2) 0.14
- (3) 0.25
- (4) 0.14
- (5) 0.14
- (6) 0.12
- (7) 0.08 and
- (8) 0.11

30. (currently amended) A population of variegated DNA sequences that encode a kappa light chain CDR1 selected from the group consisting of:

- (1) RASQ<1>V<2><2><3>LA (SEQ ID NO: 14)
- (2) RASQ<1>V<2><2><2><3>LA (SEQ ID NO: 15);

wherein <1> is an equimolar mixture of amino acid residues ADEFGHIKLMNPQRSTVWY; <2> is 0.2 S and 0.044 of each of ADEFGHIKLMNPQRTVWY; and <3> is 0.2Y and 0.044 each of ADEFGHIKLMNPQRTVW and Y; and

(3) mixtures of variegated DNA sequences
characterized by any of the above DNA sequences.

31. (original) The population of variegated DNA
sequences of claim 30, wherein CDR1s (1) and (2) are
present in the population in a ratio of 0.68:0.32.

32. (original) A population of variegated DNA
sequences that encode a kappa light chain CDR2 having the
sequence:

<1>AS<2>R<4><1>,

wherein <1> is an equimolar mixture of amino acid
residues ADEFGHIKLMNPQRSTVWY; <2> is 0.2 S and 0.044 of
each of ADEFGHIKLMNPQRTVWY; and <4> is 0.2 A and)0.044
each of DEFGHIKLMNPQRSTVWY.

33. (currently amended) A population of
variegated DNA sequences that encode a kappa light chain
CDR3 selected from the groups consisting of:

(1) QQ<3><1><1><1>P<1>T (SEQ ID NO: 16),

wherein <1> is an equimolar mixture of amino
acid residues ADEFGHIKLMNPQRSTVWY; <3> is 0.2 Y and 0.044
each of ADEFGHIKLMNPQRTVW;

(2) QQ33111P, wherein 1 and 3 are as defined
in (1) above;

(3) QQ3211PP1T (SEQ ID NO: 17), wherein 1 and 3 are as defined in (1) above and 2 is 0.2 S and 0.044 each of ADEFGHIKLMNPQRTVWY; and

(4) mixtures of variegated DNA sequences characterized by any of the above DNA sequences.

34. (original) The population of variegated DNA sequences according to claim 33, wherein CDR3s (1), (2) and (3) are present in the population in a ratio of 0.65:0.1:0.25.

35. (currently amended) A population of variegated DNA sequences that encode a lambda light chain CDR1 selected from the group consisting of:

(1) TG<1>SS<2>VG<1><3><2><3>VS (SEQ ID NO: 18),

wherein <1> is 0.27 T, 0.27 G and 0.027 each of ADEFGHIKLMNPQRSVWY, <2> is 0.27 D, 0.27 N and 0.027 each of ADEFGHIKLMPQIRSTVWY, and <3> is 0.36 Y and 0.036 each of ADEFGHIKLMNPQRSTVW;

(2) G<2><4>L<4><4><3><4><4>,

wherein <2> is as defined in (1) above and <4> is an equimolar mixture of amino acid residues ADEFGHIKLMNPQRSTVWY; and

(3) mixtures of variegated DNA sequences
characterized by any of the above DNA sequences.

36. (original) The population of variegated DNA
sequences according to claim 35, where CDR1s (1) and (2)
are present in the population in a ratio of 0.67:0.33.

37. (original) A population of variegated DNA
sequences that encode a lambda light chain CDR2 has the
sequence:

<4><4><4><2>RPS,

wherein <2> is 0.27 D, 0.27 N, and 0.027 each of
AEFGHIKLMPQSTVWY and <4> is an equimolar mixture of
amino acid residues ADEFGHIKLMNPQSTVW.

38. (currently amended) A population of variegated DNA
sequences that encode a lambda light chain CDR3 selected from
the group consisting of:

(1) <4><5><4><2><4>S<4><4><4><4>V,

wherein <2> is 0.27 D, 0.27 N, and 0.027 each
of AEFGHIKLMPQSTVWY; <4> is an equimolar mixture of
amino acid residues ADEFGHIKLMNPQSTVW; and <5> is 0.36 S
and 0.0355 each of ADEFGHIKLMNPQRTVWY;

(2) <5>SY<1><5>S<5><1><4>V (SEQ ID NO: 19),

wherein <1> is an equimolar mixture of
AEFGHIKLMNPQSTVWY; and <4> and <5> are as defined in
(1) above; and

(3) mixtures of variegated DNA sequence
characterized by any of the above DNA sequences.

39. (original) The population of variegated DNA
sequences according to claim 38, wherein CDR3s (1) and
(2) are present in the population in an equimolar
mixture.

40. (original) The population of variegated DNA
sequences according to claim 22 or 23 further comprising
variegated DNA sequences that encode a heavy chain CDR
selected from the group consisting of:

(1) one or more of the heavy chain CDR2s
defined in claim 24 or 25;

(2) one or more of the heavy chain CDR3s
defined in claims 26, 27, 28 or 29; and

(3) mixtures of variegated DNA sequence
characterized by (1) and (2).

41. (original) The population of variegated DNA
sequences according to claim 24 further comprising
variegated DNA sequences that encodes one or more heavy
chain CDR3s selected from the group defined in claims 26,
27, 28 or 29.

42. (original) The population of variegated DNA
sequences according to claim 40 or 41 further comprising

variegated DNA sequences that encodes a light chain CDR
selected from the group consisting of

- (1) one or more the kappa light chain CDR1s
defined in claim 30 or 31;
- (2) the kappa light chain CDR2 defined in
claim 32;
- (3) one or more of the kappa light chain CDR3s
defined in claim 33 or 34;
- (4) one or more of the kappa light chain CDR1s
defined in claim 35 or 36;
- (5) the lambda light chain CDR2 defined in
claim 37;
- (6) one or more of the lambda light chain
CDR3s defined in claim 38 or 39; and
- (7) mixtures of variegated DNA sequences
characterized by one or more of (1) through (6).

43. (original) A population of vectors comprising
the variegated DNA sequences of any one of claims 22-42.